

SEQUENCE LISTING

<110> Binley, Katie M  
Naylor, Stuart

<120> POLYNUCLEOTIDE CONSTRUCTS AND USES THEREOF

<130> 9192.16USWO

<140> PCT/GB99/03181  
<141> 1999-09-22

<150> PCT/GB98/02885  
<151> 1998-09-23

<150> GB 9901906.9  
<151> 1999-01-28

<150> GB 9903538.8  
<151> 1999-02-16

<160> 32

<170> PatentIn Ver. 2.1

<210> 1  
<211> 25  
<212> DNA  
<213> Mus sp.

<400> 1  
cgcgtcggtg caggacgtga caaat

25

<210> 2  
<211> 19  
<212> DNA  
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19

<210> 3  
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<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: OBhrel

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caggacgtga cagctagccc gggctcgaga tctgcgatct gcatctcaat tagtcagcaa 120  
ccatagtccc gcccctaact ccgcccattcc cgcccctaac tccgcccagt tccgcccatt 180  
ctccgccccca tcgctgacta attttttta tttatgcaga ggccgaggcc gcctcggcct 240  
ctg

243

<210> 4  
<211> 229  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic construct

<400> 4  
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gtcggtcgagg acgtgacatc tagagaacca tcagatgtt ccagggtgcc ccaaggacct 120  
gaaatgaccc tggccttat ttgaactaac caatcagttc gcttctcgct tctgttcgct 180  
cgcttctgct ccccgagctc aataaaagag cccacaaccc ctcactcg 229

<210> 5  
<211> 225  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic construct

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acgtcctgca cgactctaga gaaccatcag atgtttccag ggtgccccaa ggacctgaaa 120  
tgaccctgtg ctttatttga actaaccaat cagttcgctt ctcgcttctg ttcgcgcgct 180  
tctgctcccc gagctcaata aaagagccca caaccctca ctcgg 225

<210> 28  
<211> 249  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic promoter

<400> 28  
gcttagatcg tgcaggacgt gacatctagt gtcgtgcagg acgtgacatc tagtgcgtg 60  
caggacgtga cagctagcat tccatcacgt ggcccgagag aagcatccgg agtactacaa 120  
ggactgctga cagcgagatt tctacaaggg actttccgct ggggactttc cagggaggtg 180  
tggcctgggc gggactgggg agtggcgagc cctcagatgc tgcataataag cagcagctgc 240  
ttttcccc 249

<210> 29  
<211> 273  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Repressed  
promoter

<400> 29  
gctagagtcg tgcaggacgt gacatctagt gtcgtgcagg acgtgacatc tagtgcgtg 60  
caggacgtga cagctagcat tccatcacgt ggcccgagag aagcatccgg agtactacaa 120  
ggactgctga cagcgagatt tctacaaggg actttccgct ggggactttc cagggagggtg 180  
tggcctgggc gggactgggg agtggcaagt gaaagtgaaa gtgaaagtga gagccctcag 240  
atgctgcata taagcagcag ctgctttgc ccc 273